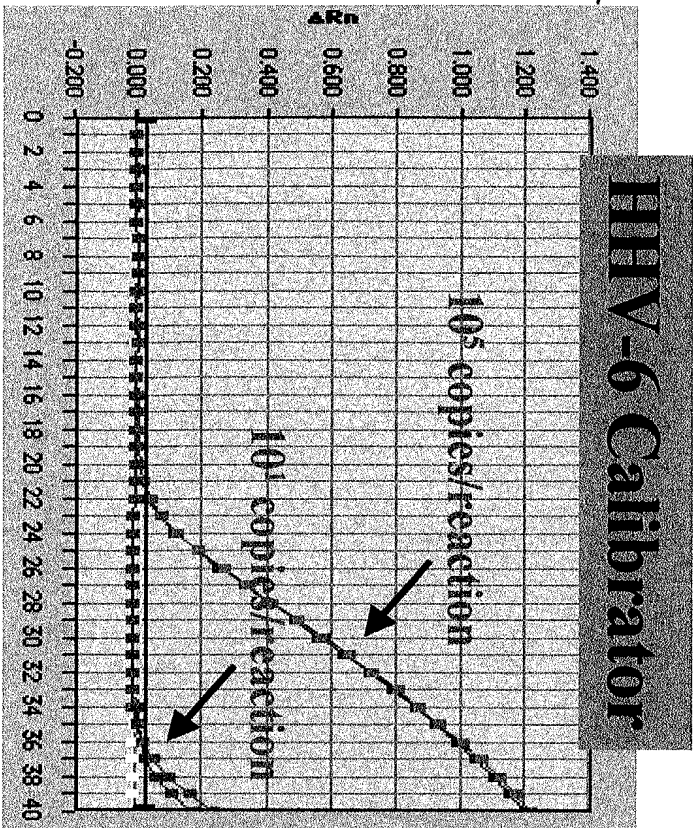
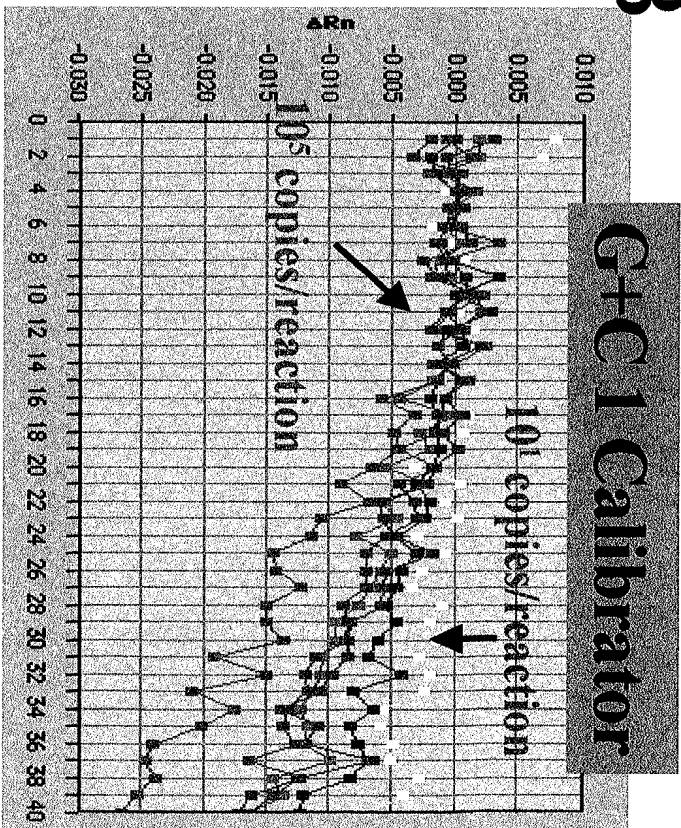


Rif^r and S^r photo

A



B



C

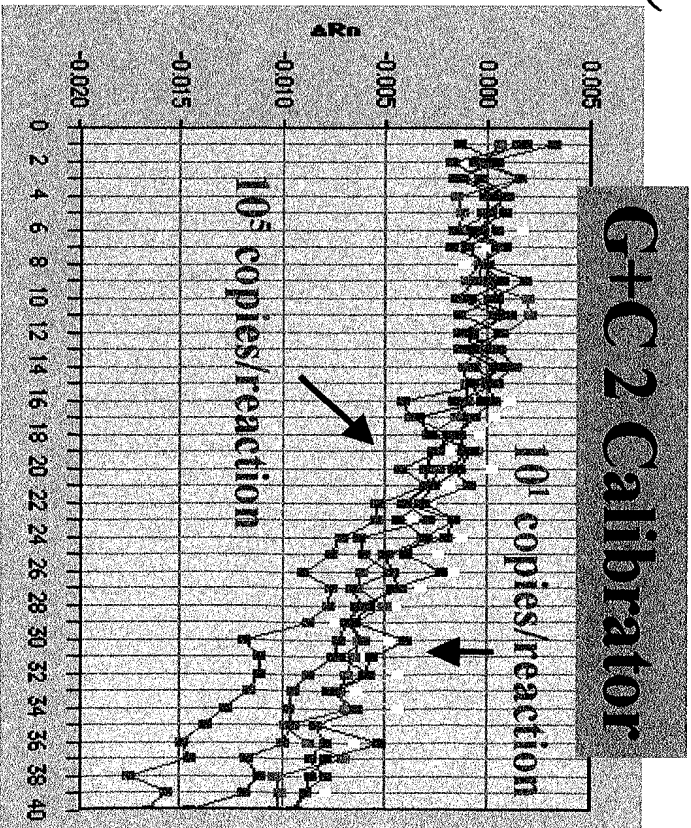


Table 1

HHV-7	Primers	Probes
	Forward (SEQ ID NO:1) 5' AGCGGTACCTGTAAAATCATCCA3' Reverse (SEQ ID NO:2) 5' AACAGAAACGCCACCTCGAT 3'	Standard (SEQ ID NO:3) 5' ACCAGTGAGAACATCGCTCTAACTGGATCA 3' Calibrator (SEQ ID NO:4) 5' TAAGCCCTGACCCGCACGGGTATAATACTAA 3'
HHV-8	Primers	Probes
	Forward (SEQ ID NO:5) 5' GTCCAGACGATATGTGCCG3'	Standard (SEQ ID NO:7) 5' CATTGGTGGTATATAGATCAAGTTCCGCCA3'
	Reverse (SEQ ID NO:6) 5' ACTCCAAAATATCGGCCGG3'	Calibrator (SEQ ID NO:8) 5' ACTATTCCATGCCGAATTGAGCATAGTTG3'

(continued)

Table 1 (continued)

	Primers	Probes
HIV-1	Forward (SEQ ID NO:9)	Standard (SEQ ID NO:11)
	5' TACTGACGCTCTCGCACC 3'	5' ATCTCTCTCCTTCTAGCCTCCGCTAGTCAA 3'
	Reverse (SEQ ID NO:10)	Calibrator (SEQ ID NO:12)
	5' TCTCGACGCGAGGACTCG 3'	5' ACTCTCAGCGGCATTCTCCTCACTTCTACT 3'
	Primers	Probes
CAMV	Forward (SEQ ID NO:13)	Standard (SEQ ID NO:15)
	5' GTCTTGCGAAGGATAGTGGGA 3'	5' TGGCTCATCCCTTACGTCAGTGAGAT 3'
	Reverse (SEQ ID NO:14)	Calibrator (SEQ ID NO:16)
	5' CACGTCTTCAAGCAAGTGGA 3'	5' ATCGCTACATGCTAAGGCATCTGTGTGC 3'

(continued)

Table 1 (continued)

	Primers	Probes
Myc.	Forward (SEQ ID NO:17)	Standard (SEQ ID NO:19)
T. 1	5' AGGAGGAGTGGCGCTGATG 3'	5' ACGAGGAGTCGCTGGCCGATCC 3'
	Reverse (SEQ ID NO:18)	Calibrator (SEQ ID NO:20)
	5' ACTCGGCGAGAGCTGCC 3'	5' TCCAGCGTCAGGCGTAGGCAGC 3'
	Primers	Probes
Myc.	Forward (SEQ ID NO:21)	Standard (SEQ ID NO:23)
T. 2	5' AGGCGAACCCTGCCCAG 3'	5' TCGACACATAGGTGAGGTCTGCTACCCACA 3'
	Reverse (SEQ ID NO:22)	Calibrator (SEQ ID NO:24)
	5' GATCGCTGATCCGCCA 3'	5' ACTACGACTACGGCTGCGATCGACATCGAT 3'

Table 2

Primers		Probes
HHV-6	Forward (SEQ ID NO:25)	Standard (SEQ ID NO:27)
	5' CAAAGCCAAATTATCCAGAGCG 3'	5' CACCAGACGTCACACCCGAAGAAT 3'
	Reverse (SEQ ID NO:26)	Calibrator (SEQ ID NO:28)
	5' CGCTAGGTTGAGGATGATCGA 3'	5' TACGCAACGCCAACAGACCTAGCGA 3'
	Calibrator	
	Primer forward (SEQ ID NO:29)	Probe (SEQ ID NO:31)
	5' CCGGAAACCGAACATTACTGAA 3'	5' TACGCAACGCCAACAGACCTAGCGA 3'
	Primer reverse (SEQ ID NO:30)	
	5' TTACGTGAGGATGATCGAGGC 3'	

Table 1

	Primers	Probes
HHV-7	Forward <u>(SEQ ID NO:1)</u>	Standard <u>(SEQ ID NO:3)</u>
	5'AGCGGTACCTGTAAAATCATCCA3'	5' ACCAGTGAGAACATCGCTCTAACTGGATCA 3'
	Reverse <u>(SEQ ID NO:2)</u>	Calibrator <u>(SEQ ID NO:4)</u>
	5' AACAGAAACGCCACCTCGAT 3'	5' TAAGCCCTGACCGCACGGGTATAATACTAA 3'
HHV-8	Primers	Probes
	Forward <u>(SEQ ID NO:5)</u>	Standard <u>(SEQ ID NO:7)</u>
	5'GTCCAGACGATATGTGCGC3'	5'CATTGGTGTATATAGATCAAGTTCCGCCA3'
	Reverse <u>(SEQ ID NO:6)</u>	Calibrator <u>(SEQ ID NO:8)</u>
	5'ACTCCAAAATATCGCGCGG3'	5'ACTATTCCATGCGGAATTCGAGCATAGTTG3'

(continued)

Table 1 (continued)

HIV-1	Primers	Probes
	Forward <u>(SEQ ID NO:9)</u> 5' TACTGACGCTCTCGCACC 3'	Standard <u>(SEQ ID NO:11)</u> 5' ATCTCTCTCCTTCTAGCCTCCGCTAGTCAA 3'
	Reverse <u>(SEQ ID NO:10)</u> 5' TCTCGACGCGAGGACTCG 3'	Calibrator <u>(SEQ ID NO:12)</u> 5' ACTCTCAGCGGCGATTCTCCTCACTTCTACT 3'
	Primers	Probes
CAMV	Forward <u>(SEQ ID NO:13)</u> 5' GTCTTGCGAAGGATAGTGGGA 3'	Standard <u>(SEQ ID NO:15)</u> 5' TGGCTCATCCCTTACGTCAGTGAGAT 3'
	Reverse <u>(SEQ ID NO:14)</u> 5' CACGTCTTCAAGCAAGTGGA 3'	Calibrator <u>(SEQ ID NO:16)</u> 5' ATCGCTACATGCTAGGCATCTGTGTGC 3'

(continued)

Table 1 (continued)

	Primers	Probes
Myc.	Forward (<u>see ID no: 17</u>) 5' AGGAGGAGTGGCGCTGATG 3'	Standard (<u>see ID no: 19</u>) 5' ACGAGGAGTCGCTGGCCGATCC 3'
T. 1	Reverse (<u>see ID no: 18</u>) 5' ACTCGGCGAGAGCTGCC 3'	Calibrator (<u>see ID no: 20</u>) 5' TCCAGCGTCAGGCGTAGGCAGC 3'
	Primers	Probes
Myc.	Forward (<u>see ID no: 21</u>) 5' AGGCGAACCCTGCCCAG 3'	Standard (<u>see ID no: 23</u>) 5' TCGACACATAGGTGAGGTCTGCTACCCACA 3'
T. 2	Reverse (<u>see ID no: 22</u>) 5' GATCGCTGATCCGCCA 3'	Calibrator (<u>see ID no: 24</u>) 5' ACTACGACTACGGCTGGATCGACATCGAT 3'

Table 2

Primers	Probes
Forward (<u>SEQ ID NO: 25</u>)	Standard (<u>SEQ ID NO: 27</u>)
5' CAAAGCCCAATTATCCAGAGCG 3'	5' CACCAGACGTCACACCCGGAAGGAAT 3'
Reverse (<u>SEQ ID NO: 26</u>)	Calibrator (<u>SEQ ID NO: 28</u>)
5' CGCTAGGTTGAGGATGATCGA 3'	5' TACGCAACGCCAACAGACCTAGCGA 3'
HHV-6	
Calibrator	
Primer forward (<u>SEQ ID NO: 29</u>)	Probe (<u>SEQ ID NO: 31</u>)
5' CCGGAAACCGAACATTACTGAA 3'	5' TACGCAACGCCAACAGACCTAGCGA 3'
Primer reverse (<u>SEQ ID NO: 30</u>)	
5' TTACGTGAGGATGATCGAGGC 3'	